

TABLE 1

Target ALSV subgroups and oligonucleotide primers used in RT-PCR assay

Target Subgroup	Primer	Sequence (5'---3') ^a	Orientation	Position ^b	Product
A	PA1	CTACAGCTGTTAGGTTCCCACT	Forward SEQ ID No: 7	gp85 of env (5564-5586)	1600/2900
	PA2	GTCACCACTGTCGCCATCCG	Reverse SEQ ID No: 8	gp85 of env (5772-5791)	
A,B,C,D, E	PU1	CTRCARCTGYTAGGYTCCCAGT	Forward SEQ ID No: 15	gp85 of env (5564-5585)	229
	PU2	GYCAYCACTGTCGCCRTCCG	Reverse SEQ ID No: 16	gp85 of env (5772-5791)	
A	PA10	GGCTTCAGGCCAAAAGGGGT	Forward SEQ ID No: 17	hr1 of gp85 (5642-5661)	232
	PA20	GTGCATTGCCACAGCGGTACTG	Reverse SEQ ID No: 18	hr2 of gp85 (5858-5879)	
B	PB1	GGCTTTACCCCATACTGATAG	Forward SEQ ID No: 19	hr1 of gp85 (5642-5661)	259
	PB2	ACACATCCTGACAGATGGACC A	Reverse SEQ ID No: 20	hr2 of gp85 (5861-5882)	
C	PC1	TATTCGCCCCAACAGGGCAC	Forward SEQ ID No: 21	hr1 of gp85 (5642-5661)	238
	PC2	CCACGTCTCCACAGCGGTAAAG T	Reverse SEQ ID No: 22	hr1 of gp85 (5858-5879)	
D	PD1	GGCTTCACCCATACGGCAG	Forward SEQ ID No: 23	hr1 of gp85 (5642-5661)	258
	PD2	CCATACGTCCACAGATAGA ATA	Reverse SEQ ID No: 24	hr1 of gp85 (5858-5882)	
E	PE1	GGCTTCGCCCCACACTCCAA	Forward SEQ ID No: 25	hr1 of gp85 (5642-5661)	265
	PE2	GCACATCTCCACAGGTGTAAAT	Reverse SEQ ID No: 26	hr1 of gp85 (5867-5888)	

^aR=A/G; Y=T/C. ^bPosition of primers corresponds to the numbering of the RNA

genome of the Prague C strain of RSV (Schwartz et al., *Cell*, 32:853-869 1983).

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